

REPLACEMENT SHEET

+1 TCACTAGTACCATATAATTACAAACACACACACACCTCAGAAACGAAGCTTGC
+53 ACATAATGTCTCTCAACAAGTACCCACAACAGAAACAACCACAGGAAGAA
1 M S L Q Q V P T T E T T T R K N
+105 CCGAAACGAGGGCGGAAACGATTACCGACAAACAAATAAGTTCCCTAGAG
17 R N E G R K R F T D K Q I S F L E
+157 TACATGTTGAGACACAGTCGAGACCCGAGTTAAGGATGAAACACCAGTTGG
34 Y M F E T Q S R P E L R M K H Q L
+209 CACATAAAACTCGGGCTTCATCTCGTCAAGTGGCGATATGGTTCCAGAACAA
51 A H K L G L H P R Q V A I W F Q N N K
+261 ACGCGCGCAGTCAAAGTCGAGGCAGATTGAGCAAGAGTATAACCGCTAAAG
69 R A R S K S R Q I E Q E Y N A L K
+313 CATAACTACGAGACGCTTGCGTCTAAATCCGAGTCTCTAAAGAAAGAGAATC
86 H N Y E T L A S K S E S L K K E N
+365 AGGCCCTACTCAATCAGgtatggttcaaacttacaatgtgcattcaacta
103 Q A L L N Q
+417 tttaagtagtttgaaattttgtgacaataaagattgacaaatgttggta
+469 taattgattaacagtGGAGGTGCTGAGAAATGTAGCAGAAAAGCATCAAGA
109 L E V L R N V A E K H Q E
+521 GAAAACTAGTAGTAGTGGCAGCGGTGAAGAATCGGATGATCGGTTACGAAC
122 K T S S S G S G E E S D D R F T N
+573 TCTCCGGACGTTATGGTCAAGAAATGAATGTTCCGTTTGGACGGTT
139 S P D V M F G Q E M N V P F C D G
+625 TTGCGTACTTTGAAGAAGGAAACAGTTGGAGATTGAAGAACAACTGCC
156 F A Y F E E G N S L L E I E E Q L P
+677 AGACCCTCAAAAGTGGGGAGTTCTAAAGAGTAAAGAAGGATGTAGAAGTA
174 D P Q K W W E F *
+729 GTAGAGTAAAAACTAAACATACCAAGATAGTTGGTTACACTTG

Athb-1 LPEKKRRLTTEOVHLLEKSFETENKLEPERKTQIAKKLGLOPQRAVWFQNRARWKTQLERDYDLKSTYDOLLSNYDSIVMDNDKLRSEVTSLTEK
Athb-6 LSEKKRRLSINQVKALEKNELENKLEPERKVLAQELGLQPRQVAVWFQNRARWKTQLEKDYGVLKTOQYDSLHRHNFDSIIRRDNESLQBISKLTK
Athb-7 NKNNQRRFSDEQIKSLEMMFESETRLEPRKKVQARELGLQPRQVAIWFKQNKRARWKSQLETEYNILRQNYDNLASOPESLKKEKQALVSELQRILKEA
Athb-12 KSNNQRRFNEEOIKSLELIFESETRLEPRKKVQARELGLQPRMTIWFKQNKRARWKTQLEKEYNTLRANYNNLASQFEIMKEKOSLVSELQRINEE
Hahb-4 RNEGGRKFDTDKOISFLEYMFETQSRPELRMKHQLAHKLGHPRQVAIWFKQNKRARSKSRQIEQYNAKHNYESLASKSESLLKENQALLENQIEVLRNV

FIG. 1

REPLACEMENT SHEET

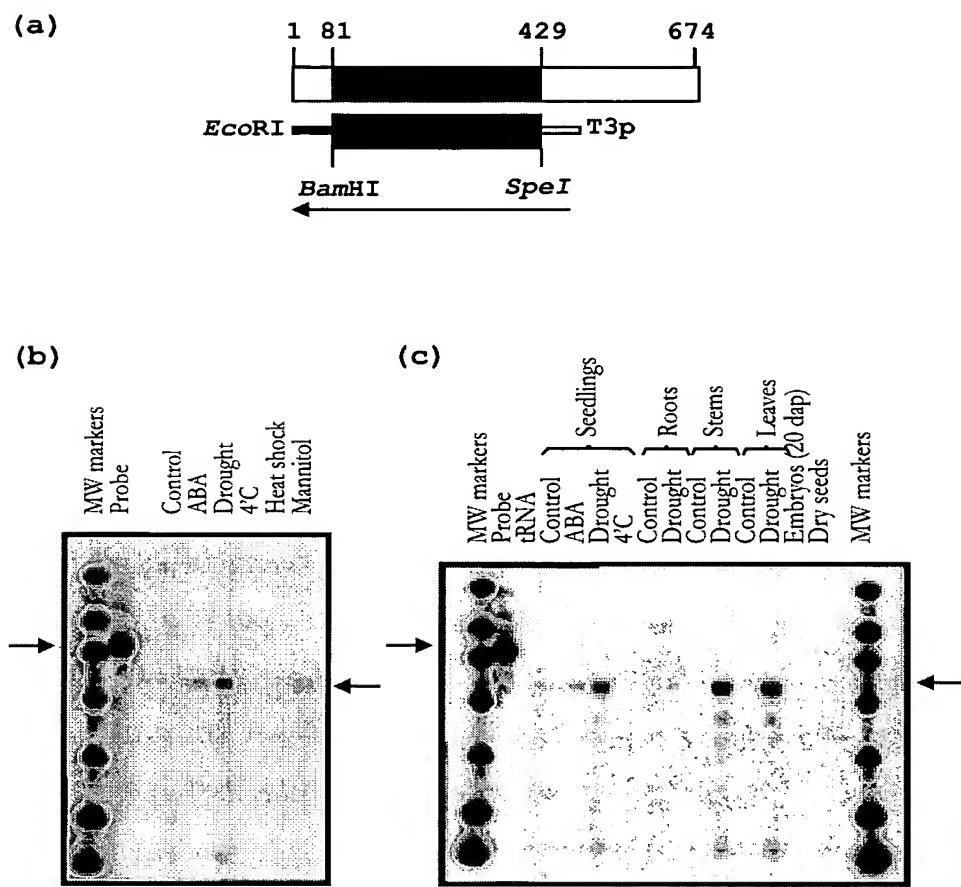


FIG. 2

REPLACEMENT SHEET

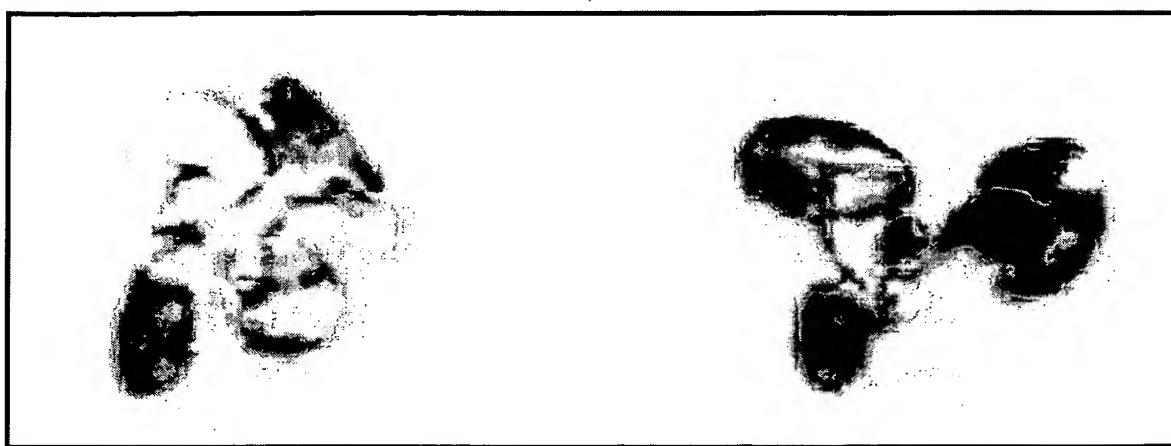


FIG. 22

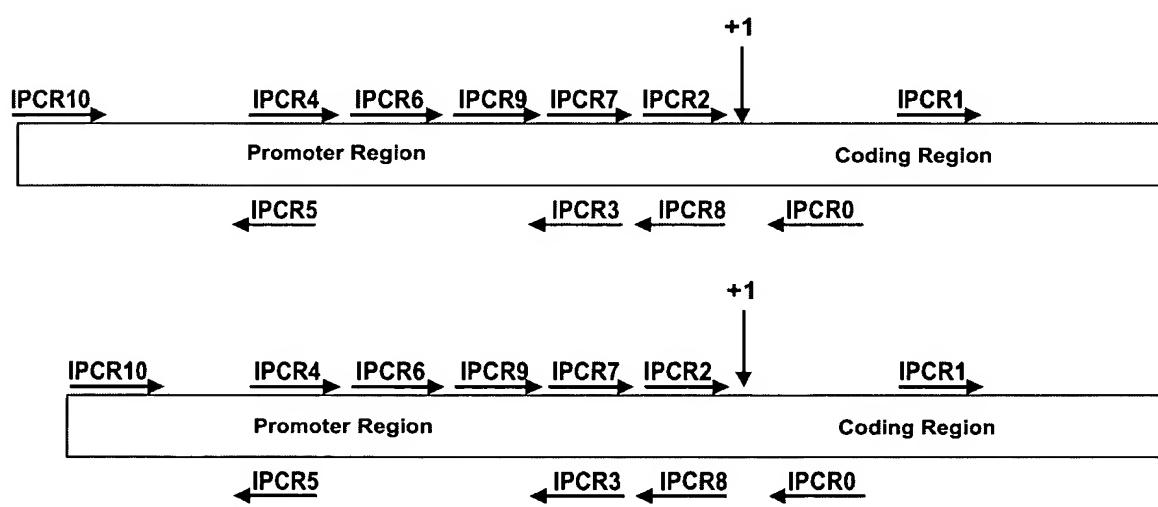


FIG. 23